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# Kisspeptin-10 Enhanced Egg Production in Quails Associated with the Increase of Triglyceride Synthesis in Liver

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ABSTRACT: Our previous results showed that kisspeptin-10 (Kp-10) injections via intraperitoneal (i.p.) once daily for three weeks notably promoted the egg laying rate in quails. In order to investigate the mechanism behind the effects of Kp-10 on enhancing the egg laying rate in birds, this study focused on the alternations of lipids synthesis in liver after Kp-10 injections. 75 female quails (22 d of age) were allocated to three groups randomly, and subjected to 0 (control, Con), 10 nmol (low dosage, L) and 100 nmol (high dosage, H) Kp-10 injections via i.p. once daily for three weeks, respectively. At d 52, quails were sacrificed and sampled for further analyses. Serum E<sub>2</sub> concentration was increased by Kp-10 injections, and reached statistical significance in H group. Serum triglyceride (TG) concentrations were increased by 46.7% in L group and 36.8% in H group, respectively, but did not reach statistical significance, and TG contents in liver were significantly elevated by Kp-10 injections in a dose-dependent manner. Serum total cholesterol (Tch) concentrations significantly decreased in H group, while in H group the hepatic Tch content was markedly increased. The level of nonesterified fatty acid (NEFA), apolipoprotein A1 and B (apoA1 and apoB) were not altered by Kp-10 injections. The genes expression of sterol regulatory element binding protein-1 (SREBP-1), fatty acid synthetase (FAS), apolipoprotein VLDL-II (apoVLDL-II), cholesterol 7α-hydroxylase (CYP7A1) and vitellogenin II (VTG-II) were significantly up-regulated by high but not low dosage of Kp-10 injection compared to the control group. However, the expression of SREBP-2, acetyl-CoA carboxylase (ACC<sub>α</sub>), malic enzyme (ME), stearoyl-CoA ( $\Delta$ 9) desaturase 1 (SCD1), apolipoprotein A1 (apoA1), fatty acid binding protein 2 (FABP2), 3-hydroxyl-3-methyl glutarylcoenzyme A reductases (HMGCR), estrogen receptor  $\alpha$ ,  $\beta$  (ER $\alpha$  and  $\beta$ ) mRNA were not affected by Kp-10 treatment. In line with hepatic mRNA abundance, hepatic SREBP1 protein content was significantly higher in H group. Although the mRNA expression was not altered, the content of ERα protein in liver was also significantly increased in H group. However, SREBP-2 protein content in liver was not changed by Kp-10 treatment. In conclusion, exogenous Kp-10 consecutive injections during juvenile stage significantly advanced the tempo of egg laying in quails, which was associated with the significant elevation in hepatic lipids synthesis and transport. (Key Words: Kisspeptin-10, Egg Laying, Triglyceride, Liver, Quail)

# INTRODUCTION

It's well documented that kisspeptin/G protein-coupled receptor 54 (GPR54) signal system is a pivotal signal pathway to regulate gonadotrophin releasing hormone (GnRH) secreting from hypothalamus and then to control the onset of normal puberty (Seminara et al., 2003). The importance of kisspeptin in normal reproductive maturation has been demonstrated by the observations that mutations in either the *Kiss-1* gene or its cognate receptor *Kiss-1R* gene render the animal unable to reach puberty (de Roux., 2003; d'Anglemont et al., 2007). In contrast, injection of kisspeptin causes the sexual precosity, and kisspeptins are

likely the most potent elicitors of GnRH/gonadotropin secretion known so far (Roa et al., 2009). Studies on rodents and primates showed that the expression of *kiss-1* gene and protein was up-regulated in hypothalamus during reproductive puberty development (Han et al., 2005; Clarkson and Herbison, 2006; Keen et al., 2008), and further activated in response to exogenous sex steroids (Smith et al., 2005). These findings proved kisspeptin as an important internal signal to control the initiation and maintenance of reproduction.

The avian neuroendocrine and endocrine system is very similar to the mammalian system (Wingfield, 2005). As in mammals, the avian reproduction is controlled by the axis of hypothalamus-pituitary gonad (HPG) and the secretion of GnRH from hypothalamus is the key step for initiating the reproduction. Although there is no evidence for a Kiss-like

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gene in birds and this gene has been suggested to have been lost in avian species during evolution (Um et al., 2010). Saldanha et al. (2010) reported that neurons that express kisspeptin-like immunoreactive (ir) protein were observed in the medial preoptic nucleus (POM) of hypothalamus in mallard drake (Saldanha et al., 2010). Kisspeptin central administration with low doses was sufficient to increase plasma LH levels in the drake and the white-crowned sparrows (Saldanha et al., 2010; Charlier et al., 2011). Our previous also showed studies that kisspeptin immunoreactive substance is expressed in ovarian granulosa cells of hens, and kisspeptin-10 stimulated progesterone secretion in cultured hens' granulosa cells from preovulatory follicles (Xiao et al., 2011). observations suggest the highly conserved function of kisspeptins in regulating reproduction across the vertebrates. However, till now, the data about the effects of kisspeptins on the avian reproduction is really scarce.

It is generally accepted that the onset of egg laying in female poultry is the indicator of puberty complement and sex maturation, and the reproductive behavior is mainly regulated by hormones secreted from HPG axis (Gous et al., 2000). In chicken each egg yolk contains about 4 gram of triglycerides (TG) (Bujo et al., 1997), thus egg production is an energy-intensive process, requiring a large increase in TG synthesis to support the demands of new yolk formation. Birds have the ability to store large quantities of excess energy (in the form of TG) in liver, adipose tissue and in yolk of developing oocytes (Hermier, 1997). The liver is known to be the primary site for lipogenesis in birds, and most of the body's endogenous lipids are hepatic origin (Xu et al., 2003). As in mammals, enzymes in the avian liver control sterol and lipid metabolism, and the levels of mRNA expression of lipogenic genes have been reported to parallel changes in enzyme activity, such as fatty acid synthase (FAS) (Wilson et al., 1986).

Several mature peptides encoded by kiss-1 gene including kisspeptin-54, -14, -13 and -10, all of their carboxyl terminus is a motif of arginine-phenylalanine-NH2 (RF-NH2), and each can activate GPR54 with equal biopotency (Kotani et al., 2001; Oakley et al., 2009). The amino acids sequence and the function of kisspeptin-10 is highly conserved among different species, including human, rat, mouse and fish (van Aerle et al., 2008). Therefore, the aim of this study was to investigate the effect of Kp-10 consecutive injection via i.p. on the initiation of egg laying in Japanese quails. The Japanese quail has been suggested as a good model for studying reproductive aging of hypothalamic systems (Ottinger et al., 2004), and an ideal species for studies of the neuroendocrine control of reproduction and sexual behavior due to the small size and early sexual maturation. To uncover the mechanism behind

the effects of Kp-10 on the onset of egg laying in birds, this study focused on the changes of cholesterol and TG synthesis in liver and transport in peripheral system, with particular emphasis on the genes expression involved in fatty acid synthesis and cholesterol metabolism.

# **MATERIAL AND METHODS**

#### **Animals**

Seventy-five female Japanese quails were obtained from a local breeding farm (Wuxi, China) and fed following the standard procedure. Quails were allowed to water *ad libitum* during the whole experimental stage. At 7 d of age, quails were randomly allocated to 15 cages (5 quails in each cage) and raised in the same place and management as well as the same diet. At 21 d of age, 15 cages were randomly divided to three groups (5 cages for each group) and subjected to 300 µl saline injection with or without Kp-10 (Cat. No. 2570, TOCRIS, USA). 300 µl saline without Kp-10 (control, Con), low (10 nmol, L) or high (100 nmol, H) Kp-10 was injected to juvenile quails via i.p. once daily for 3 wks. The first egg was observed at 42 d of age (d 42), and the egg laying performance was recorded daily till d 52.

At d 52, quails were sacrificed by cervical dislocation for samples collection and analysis. Serum was collected and stored at  $-20^{\circ}$ C, and liver tissues were removed and weighted, and quickly snap-frozen in liquid nitrogen. Frozen tissues were stored at  $-70^{\circ}$ C prior to gene and protein expression analyses. Fifteen frozen tissues were selected randomly (3 samples from each cage) and used for gene and protein expression analyses (n = 15).

Animal care and use was approved by the Institutional Animal Care and Use Committee of Nanjing Agricultural University.

#### **Materials**

Kisspeptin-10 (Cat. No. 2570) was purchased from Tocris Bioscience, USA. The contents of TG, Tch, NEFA, apoA1 and apoB were measured by the commercial kits purchased from Beijing Applygen Company (Beijing, China). Moloney Murine Leukemia Virus reverse transcriptase (M-MLV) and RNase inhibitor were products of Promega, USA. Random hexamer primers and dNTP were purchased from TaKaRa, Japan. SYBR Green Realtime PCR Master Mix was product of TOYOBO Ltd., Japan.

#### Radioimmunoassay for serum E2

The serum  $E_2$  was measured with radioimmunoassay (RIA) using commercial kits purchased from Shanghai Institute of Biological Products. The ranges of assay sensitivity for  $E_2$  were between 5 and 4,000 pg/ml. The inter- and intra-assay coefficient of variations were 10%

and 15%, respectively.

# Gene expression analyses

Total RNA was extracted from liver tissues using TRIZOL reagent (Takara, Japan). The RNA concentration and quality was then quantified by measuring the absorbance at 260 nm in a photometer (Eppendorf Biophotometer, Germany). Ratios of absorption (260/280 nm) of all preparations were between 1.8 and 2.0. Aliquots of RNA samples were subjected to electrophoresis with 1.4% agarose-formaldehyde gels stained with ethidium bromide to verify their integrity.

Reverse transcription was performed using the total RNA (2 μg) described above in a final volume of 25 μl containing 1×RT-buffer, 100 U Moloney Murine Leukemia Virus reverse transcriptase (M-MLV) (Promega, USA), 8 U RNase inhibitor (Promega, USA), 5.3 μM random hexamer primers and 0.8 mM dNTP (TaKaRa, Japan). After incubation at 37°C for 1 h, the reaction was terminated by heating at 95°C for 5 min and quickly cooling on ice.

Real-time PCR was performed in Mx3000P (Stratagene, USA). Mock RT and No Template Controls (NTC) were set to monitor the possible contamination of genomic DNA both at RT and PCR. The standard curves for each target gene, and melting curves were performed to insure a single specific PCR product for each gene. Two microliter of 16fold dilution of RT product was used for PCR in a final volume of 25 µl containing 12.5 µl SYBR Green Real-time PCR Master Mix (TOYOBO Ltd., Japan) and 0.6 to 1.0 µM of each forward and reverse primers for target genes and internal standard gene (the information of primers was shown in Table 1). Chicken β-actin mRNA was used as a reference gene for normalization purpose. The following PCR protocols were initial denaturation (1 min at 95°C), then a three-step amplification program (20 s at 95°C, 20 to 30 s at 60 to 64°C, 20 s at 72°C) was repeated 45 times. The PCR products for each gene were sent to Haojia Biotech, Ltd., for sequencing to verify the specificity. The reported sequences exactly matched those published in GenBank. The method of  $2^{-\Delta\Delta Ct}$  was used to analyze the real-time RT-

Table 1. Primer sequences of the target genes

Target genes GenBank accession number		Primer sequences (F: forward, R: reverse)	PCR products (bp)	Annealing temperature (°C)	
β-actin	NM205518	F: 5'-TGCGTGACATCAAGGAGAAG-3' R: 5'-TGCCAGGGTACATTGTGGTA -3'	300	62	
SREBP-1	AY029924	F: 5'-CTACCGCTCATCCATCAACG-3' R: 5'5 \section -CTGCTTCAGCTTCTGGTTGC-3'	145	64	
FAS	J04485	F: 5'-TGAAGGACCTTATCGCATTGC-3' R: 5'-GCATGGGAAGCATTTTGTTGT -3'	195	62	
ΑССα	NM-205505	F: 5'-AGTCCTGATTGAGCATGGCA-3' R: 5'-CTCCAGATGGCGGTAGATTC-3'	127	62	
ME	AF408407	F: 5'-AGTGCCTACCTGTGATGTTG-3' R: 5'-GGCTTGACCTCTGATTCTCT-3'	101	62	
SCD1	NM_204890	F: 5'-AGTGGTGTTGCTGTGCTTCA-3' R: 5'-CTAAGGTGTAGCGCAGGATG-3'	107	64	
HMGCR	AB109635	F: 5'- CTGGGTTTGGTTCTTGTTCA -3' R: 5'-ATTCGGTCTCTGCTTGTTCA -3'	244	64	
CYP7A1	AB109636	F: 5'-CATTCTGTTGCCAGGTGATGTT-3' R: 5'-GCTCTCTCTGTTTCCCGCTTT-3'	106	64	
SREBP-2	XM_416222	F: 5'-CCCAGAACAGCAAGCAAGG-3' R: 5'-GCGAGGACAGGAAAGAGAGTG-3'	108	64	
ApoVLDL-II	M25774	F: 5'-ATGGTGCAATACAGGGCATT-3' R: 5'-GGGAAACATCCAGCAAGAAC-3'	196	64	
FABP2	AF380999	F: 5'-GAGCTCCAGTCCCATGAAAA-3' R: 5'-TCAGCAGCTCCATCTCACAC-3'	202	64	
ApoA1	M17961	F: 5'-ATGCCATCGCCCAGTTCG-3' R: 5'-GAGCCTCGGTGTCCTTCA-3'	162	64	
VTG-II	M1806	F:5'-GCACAAGTGAAGCTGGAGTGG-3' R: 5'-AATTCTCGAGCACGGCAGAGG-3'	328	62	
ERα	X03805	F: 5'-GATATTGATGATCGGCTTAG-3' R: 5'-GTGCTCCATTCCTTTGTT-3'	418	62	
ERβ	AB036415	F:5'-GGCAGACAAAGAGCTGGTTCA-3' R: 5'-TCATGCTCAGCAGATGCTCC-3'	529	62	

PCR data. All samples were included in the same run of RT-PCR and repeated in triplicates.

# Western blot analysis

Liver tissues were immersed in 1,000 µl RIPA buffer per 200 mg. Homogenated on ice, and then kept antigraded and standed alternately for 30 min. RIPA buffer contained 50 mM Tris base (pH 7.4), 150 mM NaCl, 1 mM EDTA, 1% (v/v) NP-40, 0.25% (w/v) Na-deoxycholate, 1 mM NaF, and 1% (v/v) Halt protease inhibitor cocktail (Pierce Chemical Co., Rockford, IL, USA). Tissue lysates were centrifuged at 12,000 rpm for 10 min and the supernatant was removed. The pellet was dissolved in lysed buffer and the protein concentration was determined with the BCA (bicinchoninic acid) Protein Assay Kit (Pierce Chemical Corp., Rockford, USA). Samples were all denatured at 100°C for 5 min before loaded into a 12% polyacrylamide gel, and then transferred onto a polyvinylidene difluoride membrane (Immobilon-P, Millipore Corp., Fisher Scientific, Hanover Park, IL, USA) at 100 voltages for 90 min. The membrane was then blocked with 3% (m/v) BSA for 2 h. After that, the membranes were incubated with primary antibodies (SREBP-1 at 1:100 dilution; SREBP-2 at 1:200 dilution; ERa at 1:500 dilution and GAPDH at 1:10,000 dilution) at 4°C for 16 h. After washing, the membranes were incubated with horseradish peroxidase-conjugated anti-mouse IgG secondary antibody for 2 h at room temperature. The immunoblots were visualized by enhanced chemiluminescence detection (Pierce Chemical Corp., Rockford, USA) and quantified by integrated focal imaging system. The band densities of SRBBP-1, SREBP-2 and  $ER\alpha$  were normalized with that of GAPDH. Antibody of SREBP-1, SREBP-2 and ERα were all purchased from Abcam Company, Cambridge, UK, the catalog No. are ab3259, ab30682 and ab2746 respectively. Antibody of GAPDH was bought from Kangchen Company, Shanghai, China, and the catalog No. is KC-5G4.

#### **Statistical analysis**

The results are expressed as mean±SEM. Data were analyzed by ANOVA using the General Linear Models of SPSS 11.0 for Windows (StatSoft, Inc. USA). The effects of Kp-10 on TG and cholesterol and genes expression data were analyzed by one-way ANOVA followed by least significant difference test post hoc analysis. Data were considered statistically significant when p<0.05.

# **RESULTS**

# Serum E<sub>2</sub> concentration and lipid metabolic parameters

Serum E<sub>2</sub> concentration was markedly increased by Kp-10 injection on dose-dependent manner, and reaching the statistical significance in H group (p<0.05). The level of serum triglyceride (TG) was increased by 46.7% in L group and 36.8% in H group, respectively, but did not reach the statistical significance (p>0.05), while the content of TG in liver was significantly elevated by Kp-10 injection in a dose-dependent manner in L and H group compared to the control group (p<0.05). Compared to the control group, total cholesterol (Tch) concentration significantly decreased in H group (p<0.05), while the hepatic Tch content was markedly increased by high dosage of Kp-10 administration (p<0.05). There were no significant changes of serum or hepatic Tch content in L group (p>0.05). The levels of non-esterified fatty acid (NEFA), apolipoprotein A-I (apoA1) and apolipoproteins B (apoB) were not altered by chronic Kp-10 injection (p>0.05)(Table 2).

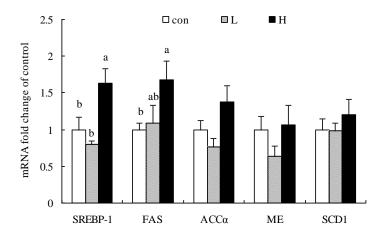
# Hepatic expression of genes involved in lipids metabolism

As shown in Figure 1 to 4, the genes expression of SREBP-1, FAS, ApoVLDL-II, CYP7A1 and VTG-II in liver were significantly up-regulated in H but not L group compared to the Con group (p<0.05). SREBP-2, ACC $_{\alpha}$ , ME, SCD1, ApoA1, FABP2, HMGCR, ER $\alpha$  and  $\beta$  mRNA

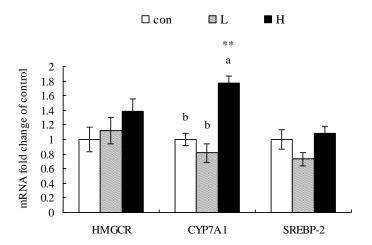
Table 2. Effects of kp-10 on the contents of parameters in serum and liver

	Con	L	Н
In serum			
Tch (mg/ml)	$2.781\pm0.115^{a}$	2.914±0.153 <sup>a</sup>	$2.443\pm0.088^{b}$
TG (mg/ml)	18.098±3.761	26.498±4.697	24.758±3.484
NEFA (mmol/L)	3.82±0.63	3.47±0.56	3.93±0.39
apoA1 (g/L)	$0.10\pm0.03$	0.11±0.04	$0.10\pm0.05$
apoB (g/L)	$0.35\pm0.04$	$0.40\pm0.08$	0.35±0.06
E <sub>2</sub> concentration (pg/ml)	$633.15\pm68.02^{a}$	703.82±27.72 <sup>a</sup>	777.54±22.77 <sup>b</sup>
In liver			
Tch (mg/100 g)	28.07±3.81 <sup>b</sup>	$32.27\pm4.00^{b}$	$37.68\pm4.07^{a}$
TG (mg/100 g)	$13.10\pm0.31^{a}$	17.13±0.91 <sup>b</sup>	23.63±1.51 <sup>b</sup> ,**

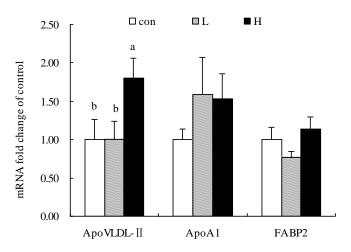
All values were mean $\pm$ SE, n = 25, Means within rows with different letter superscripts are significantly different (p<0.05), and \*\* at the level of p<0.01 (compared with control).



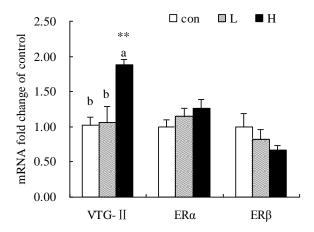
**Figure 1.** Effects of Kp-10 on the hepatic expression of lipogenic genes including SREBP-1, FAS, ACCα, ME and SCD1 mRNA in quails. Values are means±SE of fifteen replications. The data points without same superscripts (a, b) are significantly different (one-way ANOVA) at the level of p<0.05 by least square difference tests.



**Figure 2.** Effects of Kp-10 on the hepatic expression of cholesterol metabolic genes including HMGCR, CYP7A1 and SREBP-2 mRNA in quails. Values are means±SE of fifteen replications. The data points without same superscripts (a, b) are significantly different (one-way ANOVA) at the level of p<0.05 by least square difference tests.



**Figure 3.** Effects of Kp-10 on the hepatic expression of lipids transporting genes including ApoVLDL-II, ApoA1 and FABP2 mRNA in quails. Values are means±SE of fifteen replications. The data points without same superscripts (a, b) are significantly different (one-way ANOVA) at the level of p<0.05 by least square difference tests.



**Figure 4.** Effects of Kp-10 on the hepatic expression of VTG-II, ERα and ERβ mRNA in quails. Values are means±SE of fifteen replications. The data points without same superscripts (a, b) are significantly different (one-way ANOVA) at the level of p<0.05 and \*\* at the level of p<0.01 (compared with control) by least square difference tests.

expressions were not altered by Kp-10 treatment (p>0.05)(Figures 1, 2, 3 and 4).

# Correlation analyses among genes expression

Some of the significant correlations were found among genes encoding hepatic enzymes linked with lipogenic pathway. Table 3 showed that SREBP-1 gene expression

was positively correlated with FAS and ACC $\alpha$  genes (p<0.01). ACC- $\alpha$  mRNA expression in liver was positively correlated with the genes expression of SREBP-1, FAS, ME and SCD1 (p<0.05). As shown in Table 4, there was a highly positive correlation between apoVLDL-II and VTG-II genes expression in liver (p<0.01). There was a mild correlation between E2 level and VTG-II mRNA expression (p<0.1), between ER $\alpha$  and VTG-II (p<0.1), ER $\alpha$  and apoVLDL-II mRNA expression (p<0.05) (Tables 3 and 4).

# Hepatic SREBP-1, -2 and ERα protein content

Hepatic protein content of SREBP-1, 2 and ER $\alpha$  was determined with Western blot analysis and shown in Figure 5. In line with hepatic mRNA abundance, hepatic SREBP-1 content was significantly higher in H group than the control group. ER $\alpha$  Protein content in liver was also significantly increased by high dosage of Kp-10, although its mRNA expression was not altered by Kp-10 treatment. However, no obvious changes of SREBP-2 protein expression in liver were observed in quails among the Con, L and H groups (Figure 5).

#### DISCUSSION

In this study, we reported herein the effects of kisspeptin-10 on regulating the initiation of egg laying in

Table 3. Correlations between expressions of hepatic genes related to lipid synthesis

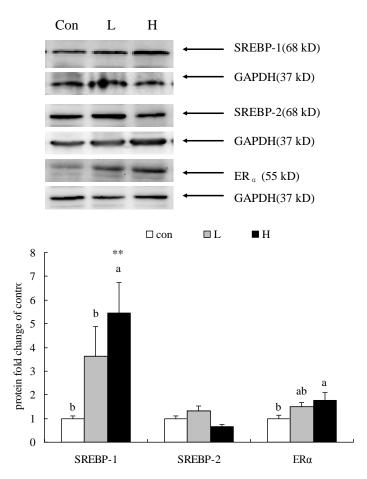
	SREBP-1	FAS	ACC-α	ME	SCD1
			r(P)		
SREBP-1	1	0.51662	0.5629	0.35621	0.28955
		(0.0049)	(0.0034)	(0.0579)	(0.1604)
FAS		1	0.50073	0.28652	0.47762
			(0.0108)	(0.1394)	(0.0158)
ACC-α			1	0.58708	0.69372
				(0.0020)	(0.0001)
ME				1	0.36621
					(0.0718)
SCD1					1

Pearson's correlation coefficients (r) are shown with p values in parentheses.

Table 4. Correlations between expressions of hepatic genes related to lipid synthesis and transporting to egg yolk

	$E_2$	ApoA1	VTG-II	ApoVLDL-II	$ER\alpha$
<del>.</del>			r(P)		
$E_2$	1	0.245	0.479	-0.045	-0.065
		(0.200)	(0.098)	(0.831)	(0.757)
ApoA1		1	-0.136	0.056	-0.040
			(0.673)	(0.797)	(0.854)
VTG-II			1	0.853	0.486
				(0.000)	(0.092)
ApoVLDL-II				1	0.489
					(0.015)
ERα					1

Pearson's correlation coefficients (r) are shown with p values in parentheses.



**Figure 5.** Effects of Kp-10 on the hepatic expression of SREBP-1, SREBP-2 and ER $\alpha$  proteins in quails. Values are means $\pm$ SE of fifteen replications. The data points without same superscripts (a, b) are significantly different (one-way ANOVA) at the level of p<0.05 and \*\* at the level of p<0.01 (compared with control) by least square difference tests.

quails for the first time. Our previous results showed that daily i.p. injection of kisspeptin-10 with 10 or 100 nmol to juvenile female quails for 3 wks notably evoked an earlier onset of egg laying (Ni et al., 2012). In this study, we found that there was a significant increase in lipids synthesis in liver, as well as the significant up-regulation of estrogen receptor protein and lipogenic genes expression in liver.

Kisspeptin encoded by *Kiss-1* gene acts as the GnRH secretagogue to play a critical role in the physiology of puberty in mammals (Navarro, 2005). The avian neuroendocrine and endocrine system is very similar to the mammalian system (Wingfield, 2005). Although there is no evidence for a *Kiss*-like gene in birds, and even this gene is suggested to be lost in avian species during evolution (Akazome et al., 2010), kisspeptin-like immunoreactivity (i.r) has been described in the central brain of Mallard drake (Saldanha et al., 2010). Intracerebroventricular injection of kisspeptin-10 into drake brain elicited a robust increase in circulating LH levels (Saldanha et al., 2010). Our results showed that consecutive Kp-10 injection significantly increased estrogen secretion and HPG axis activity (Ni et al., 2012), showing a similar biological effect of kisspeptins on

reproduction in birds as in mammals.

In all egg-laying vertebrates, there are a large amount of lipids transported to and stored in egg yolk. In chicken, each egg contains about 4 gram triglyceride (Bujo et al., 1997), thus the initiation and maintenance of egg production is an energy-intensive process and requires a large quantity of lipids (especially triglyceride) for egg yolk forming. There is fairly low lipid content in avian diets and lipogenesis in liver is the major path for lipid depositing in the adipose tissue and ovary of female poultry (Richards et al., 2003). Our results showed that paralleling with a higher rate of egg laying, Kp-10 injection significantly increased the contents of TG and Tch in livers of female quails, suggesting an improvement of hepatic lipogenesis. In the present study, the contents of Tch and TG in liver were significantly higher in H group than that in Con chickens. The increase of Tch in liver and the decrease of Tch in circulating system might contribute to transport more lipids to the follicles to form egg yolk.

In this study, high dosage of Kp-10 injections significantly increased the expression of SREBP-1c and FAS mRNA and protein synthesis of SREBP-1c in liver.

There were strong positive correlations of SREBP-1 gene expression with ACCα and FAS genes expression in liver, respectively. The up-regulation of SREBP-1c and FAS genes expression may account for the higher contents of TG in liver as well as in blood. Moreover, the expression of hepatic apoVLDL-II gene in H group was also significantly increased. ApoVLDL-II is a major apolipoprotein in plasma VLDL in laying hens and in estrogen-treated cockerels and roosters (Codina-Salada et al., 1983). Besides, ApoVLDL-II may inhibit the activities of lipoprotein lipase (LPL) (Walzem et al., 1999), then to protect the triglycerides from decomposing. It is produced exclusively in the liver in response to estrogen treatment. The mRNA expression of ApoVLDL-II increases markedly after estrogen treatment (Codina-Salada et al., 1983).

Vitellogenin (VTG), a precursor yolk protein (Tata, 1976), is synthesized in the liver under estrogenic induction (Le Menn et al., 1980; Wallace and Selman, 1985). Although the changes of VTG-II mRNA expression during different egg laying cycles have not been reported in birds, the age-related studies on the VTG-II gene of Japanese quail was presented by Gupta et al. (2006). In birds, the timing of VTG-II gene expression parallels with that of egg laying. In this study, accompanied by higher level of E2 in circulation VTG-II gene expression in liver was increased significantly in H group. In addition, there are mild correlations between E<sub>2</sub> and VTG-II mRNA expression, as well as ERa and VTG-II expressions. Positive correlations were also observed between ERα and apoVLDL-II, ApoA1 and apoB, ApoVLDL-II and VTG-II genes expression in liver.

In summary, we reported herein the first time that consecutive injections of kisspeptin-10 with 10 or 100 nmol to juvenile female quails notably evoked an earlier onset of egg laying, which paralleled with an improvement in TG and Tch synthesis in liver. The increase in hepatic lipogenic synthesis was associated with the up-regulation of hepatic SREBP-1, FAS, ApoVLDL-II and VTG-II genes expression.

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